

Naturally occurring mutations in the human 5-  
**lipoxxygenase gene** promoter that modify  
transcription factor binding and reporter gene  
transcription.

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Kemp J; Israel E; Busse W; Ledford D; Murray J J; Segal A;  
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AB Five lipoxxygenase (5-LO) is the first committed enzyme in the metabolic  
pathway leading to the synthesis of the leukotrienes. We examined genomic  
DNA isolated from 25 normal subjects and 31 patients with asthma (6 of  
whom had aspirin-sensitive asthma) for mutations in the known  
transcription factor binding regions and the protein encoding region of  
the **5-LO gene**. A family of mutations in the  
G + C-rich transcription factor binding region was identified consisting  
of the deletion of one, deletion of two, or addition of one zinc finger  
(Sp1/Egr-1) binding sites in the region 176 to 147 bp upstream from the  
ATG translation start site where there are normally 5 Sp1 binding motifs  
in tandem. Reporter gene activity directed by any of the mutant forms of  
the transcription factor binding region was significantly ( $P < 0.05$ ) less  
effective than the activity driven by the wild type transcription factor  
binding region. Electrophoretic mobility shift assays (EMSAs)  
demonstrated the capacity of wild type and mutant transcription factor  
binding regions to bind nuclear extracts from human umbilical vein  
endothelial cells (HUVECs). These data are consistent with a family of  
mutations in the **5-LO gene** that can modify  
reporter gene transcription possibly through differences in Sp1 and Egr-1  
transactivation.

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TITLE: Classifying patients with inflammatory disease, specifically  
asthma;  
disease diagnosis and estimation of suitability of therapy  
AUTHOR: Drazen J M; In K H; Asano K; Beier D; Grobholz J  
PATENT ASSIGNEE: Brigham+Women's-Hosp.Boston  
LOCATION: Boston, MA, USA.  
PATENT INFO: WO 9742347 13 Nov 1997  
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DOCUMENT TYPE: Patent  
LANGUAGE: English  
OTHER SOURCE: WPI: 1997-558997 [51]

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AB A new method for classifying patients suffering from inflammatory disease involves: identifying in DNA from at least 1 patient a sequence **polymorphism**, as compared with the normal 5-lipoxygenase (EC-1.13.11.12) gene, in a **5-lipoxygenase gene** regulatory sequence; and classifying the patient based on the identified **polymorphism**. The inflammatory disease is asthma, ulcerative colitis, sinusitis, psoriasis, allergic and non-allergic rhinitis, lupus or rheumatoid arthritis. The sequence **polymorphism** is addition or deletion of a binding site (Sp1 or Egr-1 site) for a transcriptional activator or repressor expressed in white blood cells or is a substitution or mutation which disrupts the binding site. Also new are: identification of an asthma patient who is a candidate for therapy with 5'-lipoxygenase-inhibitors by comparing the level of 5-lipoxygenase with that of a healthy person or by detecting a **5'-lipoxygenase gene** mutation; identification of a person susceptible to inflammatory disease, which involves detecting a DNA **polymorphism** in the **5-lipoxygenase gene** from the person. (56pp)

99365912 PubMed ID: 10436859

TITLE: Mutations in the human 5-lipoxygenase  
gene.  
AUTHOR: In K H; Silverman E S; Asano K; Beier D; Fischer A R; Keith  
T P; Serino K; Yandava C; De Sanctis G T; Drazen J M  
CORPORATE SOURCE: Department of Medicine, Brigham and Women's Hospital,  
Boston, MS, USA.  
SOURCE: CLINICAL REVIEWS IN ALLERGY AND IMMUNOLOGY, (1999  
Spring-Summer) 17 (1-2) 59-69.  
Journal code: 9504368. ISSN: 1080-0549.  
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AB Our data demonstrate the presence of a naturally occurring family of  
alleles in the core promoter of the **5-LO gene**  
, which is characterized by the deletion or addition of consensus Sp1  
(-GGGCGG) and Egr-1 (-GCGGGGGCG-) binding motifs. Each of the  
**variant** alleles can bind Sp1 and Egr-1 protein, as indicated by  
EMSA and supershift analysis with nuclear extracts. In addition,  
preliminary data from CAT reporter assays indicate that these alleles are  
less effective than the wild-type allele in initiating **5-  
LO gene** expression. Whether patients harboring the  
various alleles identified herein have different capacities to transcribe  
the **5-LO gene** and the importance of such  
potential regulation to the clinical expression of 5-LO have yet to be  
determined.

Genetic **polymorphisms** of 5-LO

AUTHOR(S): Silverman, Eric S.; In, Kwang H.; Collins, Tucker;  
Drazen, Jeffrey M.

CORPORATE SOURCE: Pulmonary and Critical Care Division, Department of  
Medicine, Vascular Research Division, Department of  
Pathology, Brigham and Women's Hospital and Harvard  
Medical School, Boston, MA, 02115, USA

SOURCE: Novel Inhibitors of Leukotrienes (1999), 147-164.  
Editor(s): Folco, Giancarlo; Samuelsson, Bengt;  
Murphy, Robert C. Birkhaeuser Verlag: Basel, Switz.  
CODEN: 68HHAI

DOCUMENT TYPE: Conference; General Review

LANGUAGE: English

AB A review with 43 refs. on 5-lipoxygenase (**5-LO**)  
**gene** regulation and how this regulation may be altered by  
naturally occurring promoter mutations. Topics include: regulation of  
5-LO; **5-LO gene** transcription; naturally  
occurring 5-LO promoter mutations in healthy and asthmatic humans; and Sp1  
and Egr-1 interactions in gene transcription.

Asthma pharmacogenetics

AUTHOR(S): Drazen, Jeffrey M.  
CORPORATE SOURCE: Women's Hospital, Brigham, UT, USA  
SOURCE: Pharmaceutical News (2000), 7(6), 26-31  
CODEN: PHNEEP; ISSN: 1071-894X  
PUBLISHER: G+B Magazines  
DOCUMENT TYPE: Journal; General Review  
LANGUAGE: English

AB A review with 33 refs. Topics discussed include the definition of asthma; treatments for asthma including bronchodilators, inhaled corticosteroids and anti-leukotrienes; pharmacogenetic mechanisms of asthma; repeatability of treatment responses; **polymorphism** of the .beta.2-adrenergic receptor; and **polymorphism** of the 5-lipoxygenase gene promoter.